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CORRECTION

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# Correction to: Relatively frequent switching of transcription start sites during cerebellar development

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## Correction

The authors of the original article [1] would like to recognize the critical contribution of core members of the FANTOM5 Consortium, who played the critical role of HeliScopeCAGE sequencing experiments, quality control of tag reads and processing of the raw sequencing data.

The revised author list includes authors Masayoshi Itoh, Hideya Kawaji, Timo Lassmann, Carsten O. Daub, Erik Arner, Piero Carninci, Yoshihide Hayashizaki and Alistair R. R. Forrest, as well as their affiliations.

The following updates to Author contributions, Acknowledgements and Funding have also been included below:

## New Author Contributions statement

PZ, TH, DS and DG generated samples for the time series. MI and PC generated the CAGE libraries. YH, COD, ARRF and PC managed and organized the FANTOM5 project. ED, TL, HK, EA and PZ performed data analysis. PZ performed biological validation experiments. PZ, DG, ED and WH wrote the manuscript. The authors read and approved the final manuscript.

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